RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/580,660
Source: IFW
Date Processed by STIC: 06/07/2006

ENTERED

IFWP

RAW SEQUENCE LISTING DATE: 06/07/2006
PATENT APPLICATION: **US/10/580,660** TIME: 11:06:46

Input Set : **F:\DEBE066US1.TXT**
Output Set: **N:\CRF4\06072006\J580660.raw**

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      3 <110> APPLICANT: HOFMEISTER, ROBERT ET AL.
      5 <120> TITLE OF INVENTION: Compositions comprising polypeptides
      7 <130> FILE REFERENCE: DEBE:066US
C-->  9 <140> CURRENT APPLICATION NUMBER: US/10/580,660
C--> 10 <141> CURRENT FILING DATE: 2006-05-26
     12 <150> PRIOR APPLICATION NUMBER: PCT/EP 2004/013445
     13 <151> PRIOR FILING DATE: 2004-11-26
     15 <150> PRIOR APPLICATION NUMBER: EP 03 027 511.9
     16 <151> PRIOR FILING DATE: 2003-11-28
     18 <160> NUMBER OF SEQ ID NOS: 6
     20 <170> SOFTWARE: PatentIn version 3.1
     23 <210> SEQ ID NO: 1
     24 <211> LENGTH: 504
     25 <212> TYPE: PRT
     26 <213> ORGANISM: artificial sequence
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     37             20                  25                  30
     39 Gly Asp Ser Tyr Leu Asn Trp Tyr Gln Gln Ile Pro Gly Gln Pro Pro
     40         35                  40                  45
     42 Lys Leu Leu Ile Tyr Asp Ala Ser Asn Leu Val Ser Gly Ile Pro Pro
     43     50                  55                  60
     45 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile His
     46 65                  70                  75                  80
     48 Pro Val Glu Lys Val Asp Ala Ala Thr Tyr His Cys Gln Gln Ser Thr
     49                 85                  90                  95
     51 Glu Asp Pro Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Gly
     52             100                 105                 110
     54 Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gln Val
     55         115                 120                 125
     57 Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ser Ser Val
     58     130                 135                 140
     60 Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Ser Ser Tyr Trp Met
     61 145                 150                 155                 160
     63 Asn Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile Gly Gln
     64                 165                 170                 175
     66 Ile Trp Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe Lys Gly
     67             180                 185                 190
     69 Lys Ala Thr Leu Thr Ala Asp Glu Ser Ser Ser Thr Ala Tyr Met Gln
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72 Leu Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr Phe Cys Ala Arg
73     210                 215                 220
75 Arg Glu Thr Thr Thr Val Gly Arg Tyr Tyr Tyr Ala Met Asp Tyr Trp
76 225                 230                 235                 240
78 Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Gly Ser Asp
79                 245                 250                 255
81 Ile Lys Leu Gln Gln Ser Gly Ala Glu Leu Ala Arg Pro Gly Ala Ser
82             260                 265                 270
84 Val Lys Met Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Arg Tyr Thr
85         275                 280                 285
87 Met His Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile Gly
88     290                 295                 300
90 Tyr Ile Asn Pro Ser Arg Gly Tyr Thr Asn Tyr Asn Gln Lys Phe Lys
91 305                 310                 315                 320
93 Asp Lys Ala Thr Leu Thr Thr Asp Lys Ser Ser Ser Thr Ala Tyr Met
94                 325                 330                 335
96 Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala
97             340                 345                 350
99 Arg Tyr Tyr Asp Asp His Tyr Cys Leu Asp Tyr Trp Gly Gln Gly Thr
100         355                 360                 365
102 Thr Leu Thr Val Ser Ser Val Glu Gly Gly Ser Gly Gly Ser Gly Gly
103     370                 375                 380
105 Ser Gly Gly Ser Gly Gly Val Asp Asp Ile Gln Leu Thr Gln Ser Pro
106 385                 390                 395                 400
108 Ala Ile Met Ser Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys Arg
109                 405                 410                 415
111 Ala Ser Ser Ser Val Ser Tyr Met Asn Trp Tyr Gln Gln Lys Ser Gly
112             420                 425                 430
114 Thr Ser Pro Lys Arg Trp Ile Tyr Asp Thr Ser Lys Val Ala Ser Gly
115         435                 440                 445
117 Val Pro Tyr Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu
118     450                 455                 460
120 Thr Ile Ser Ser Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln
121 465                 470                 475                 480
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131 <211> LENGTH: 505
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133 <213> ORGANISM: artificial sequence
136 <220> FEATURE:
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142 1               5                   10                  15
144 Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Ser Ser Tyr
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PATENT APPLICATION: **US/10/580,660** TIME: 11:06:46

Input Set : **F:\DEBE066US1.TXT**
Output Set: **N:\CRF4\06072006\J580660.raw**

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147 Trp Met Asn Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
148         35                  40                  45
150 Gly Gln Ile Trp Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe
151     50                  55                  60
153 Lys Gly Lys Ala Thr Leu Thr Ala Asp Glu Ser Ser Ser Thr Ala Tyr
154 65                  70                  75                  80
156 Met Gln Leu Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr Phe Cys
157                 85                  90                  95
159 Ala Arg Arg Glu Thr Thr Thr Val Gly Arg Tyr Tyr Tyr Ala Met Asp
160             100                 105                 110
162 Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Gly
163         115                 120                 125
165 Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Asp Ile Gln Leu Thr
166     130                 135                 140
168 Gln Ser Pro Ala Ser Leu Ala Val Ser Leu Gly Gln Arg Ala Thr Ile
169 145                 150                 155                 160
171 Ser Cys Lys Ala Ser Gln Ser Val Asp Tyr Asp Gly Asp Ser Tyr Leu
172                 165                 170                 175
174 Asn Trp Tyr Gln Gln Ile Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr
175             180                 185                 190
177 Asp Ala Ser Asn Leu Val Ser Gly Ile Pro Pro Arg Phe Ser Gly Ser
178         195                 200                 205
180 Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile His Pro Val Glu Lys Val
181     210                 215                 220
183 Asp Ala Ala Thr Tyr His Cys Gln Gln Ser Thr Glu Asp Pro Trp Thr
184 225                 230                 235                 240
186 Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Ser Gly Gly Gly Gly Ser
187                 245                 250                 255
189 Asp Ile Lys Leu Gln Gln Ser Gly Ala Glu Leu Ala Arg Pro Gly Ala
190             260                 265                 270
192 Ser Val Lys Met Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Arg Tyr
193         275                 280                 285
195 Thr Met His Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
196     290                 295                 300
198 Gly Tyr Ile Asn Pro Ser Arg Gly Tyr Thr Asn Tyr Asn Gln Lys Phe
199 305                 310                 315                 320
201 Lys Asp Lys Ala Thr Leu Thr Thr Asp Lys Ser Ser Ser Thr Ala Tyr
202                 325                 330                 335
204 Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
205             340                 345                 350
207 Ala Arg Tyr Tyr Asp Asp His Tyr Cys Leu Asp Tyr Trp Gly Gln Gly
208         355                 360                 365
210 Thr Thr Leu Thr Val Ser Ser Val Glu Gly Gly Ser Gly Gly Ser Gly
211     370                 375                 380
213 Gly Ser Gly Gly Ser Gly Gly Val Asp Asp Ile Gln Leu Thr Gln Ser
214 385                 390                 395                 400
216 Pro Ala Ile Met Ser Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys
217                 405                 410                 415
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Input Set : **F:\DEBE066US1.TXT**
Output Set: **N:\CRF4\06072006\J580660.raw**

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219 Arg Ala Ser Ser Ser Val Ser Tyr Met Asn Trp Tyr Gln Gln Lys Ser
220             420                 425                 430
222 Gly Thr Ser Pro Lys Arg Trp Ile Tyr Asp Thr Ser Lys Val Ala Ser
223         435                 440                 445
225 Gly Val Pro Tyr Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser
226     450                 455                 460
228 Leu Thr Ile Ser Ser Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys
229 465                 470                 475                 480
231 Gln Gln Trp Ser Ser Asn Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu
232                 485                 490                 495
234 Glu Leu Lys His His His His His His
235             500                 505
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239 <211> LENGTH: 504
240 <212> TYPE: PRT
241 <213> ORGANISM: artificial sequence
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246 <400> SEQUENCE: 3
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249 1               5                   10                  15
251 Ser Val Lys Met Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Arg Tyr
252             20                  25                  30
254 Thr Met His Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
255         35                  40                  45
257 Gly Tyr Ile Asn Pro Ser Arg Gly Tyr Thr Asn Tyr Asn Gln Lys Phe
258     50                  55                  60
260 Lys Asp Lys Ala Thr Leu Thr Thr Asp Lys Ser Ser Ser Thr Ala Tyr
261 65                  70                  75                  80
263 Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
264                 85                  90                  95
266 Ala Arg Tyr Tyr Asp Asp His Tyr Cys Leu Asp Tyr Trp Gly Gln Gly
267             100                 105                 110
269 Thr Thr Leu Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly
270         115                 120                 125
272 Ser Gly Gly Gly Gly Ser Asp Ile Gln Leu Thr Gln Ser Pro Ala Ile
273     130                 135                 140
275 Met Ser Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys Arg Ala Ser
276 145                 150                 155                 160
278 Ser Ser Val Ser Tyr Met Asn Trp Tyr Gln Gln Lys Ser Gly Thr Ser
279                 165                 170                 175
281 Pro Lys Arg Trp Ile Tyr Asp Thr Ser Lys Val Ala Ser Gly Val Pro
282             180                 185                 190
284 Tyr Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile
285         195                 200                 205
287 Ser Ser Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp
288     210                 215                 220
290 Ser Ser Asn Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys
291 225                 230                 235                 240
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PATENT APPLICATION: **US/10/580,660** TIME: 11:06:46

Input Set : **F:\DEBE066US1.TXT**
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293 Ser Gly Gly Gly Gly Ser Gln Val Gln Leu Gln Gln Ser Gly Ala Glu
294                 245                 250                 255
296 Leu Val Arg Pro Gly Ser Ser Val Lys Ile Ser Cys Lys Ala Ser Gly
297             260                 265                 270
299 Tyr Ala Phe Ser Ser Tyr Trp Met Asn Trp Val Lys Gln Arg Pro Gly
300         275                 280                 285
302 Gln Gly Leu Glu Trp Ile Gly Gln Ile Trp Pro Gly Asp Gly Asp Thr
303     290                 295                 300
305 Asn Tyr Asn Gly Lys Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Glu
306 305                 310                 315                 320
308 Ser Ser Ser Thr Ala Tyr Met Gln Leu Ser Ser Leu Ala Ser Glu Asp
309                 325                 330                 335
311 Ser Ala Val Tyr Phe Cys Ala Arg Arg Glu Thr Thr Thr Val Gly Arg
312             340                 345                 350
314 Tyr Tyr Tyr Ala Met Asp Tyr Trp Gly Gln Gly Thr Thr Val Thr Val
315         355                 360                 365
317 Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly
318     370                 375                 380
320 Ser Asp Ile Gln Leu Thr Gln Ser Pro Ala Ser Leu Ala Val Ser Leu
321 385                 390                 395                 400
323 Gly Gln Arg Ala Thr Ile Ser Cys Lys Ala Ser Gln Ser Val Asp Tyr
324                 405                 410                 415
326 Asp Gly Asp Ser Tyr Leu Asn Trp Tyr Gln Gln Ile Pro Gly Gln Pro
327             420                 425                 430
329 Pro Lys Leu Leu Ile Tyr Asp Ala Ser Asn Leu Val Ser Gly Ile Pro
330         435                 440                 445
332 Pro Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile
333     450                 455                 460
335 His Pro Val Glu Lys Val Asp Ala Ala Thr Tyr His Cys Gln Gln Ser
336 465                 470                 475                 480
338 Thr Glu Asp Pro Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
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341 Ser Gly His His His His His His
342             500
345 <210> SEQ ID NO: 4
346 <211> LENGTH: 503
347 <212> TYPE: PRT
348 <213> ORGANISM: artificial sequence
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351 <223> OTHER INFORMATION: Construct 8: VH(CD3)-VL(CD3)-VL(CD19)-VH(CD19)
353 <400> SEQUENCE: 4
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358 Ser Val Lys Met Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Arg Tyr
359             20                  25                  30
361 Thr Met His Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
362         35                  40                  45
364 Gly Tyr Ile Asn Pro Ser Arg Gly Tyr Thr Asn Tyr Asn Gln Lys Phe
365     50                  55                  60
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VERIFICATION SUMMARY DATE: 06/07/2006
PATENT APPLICATION: **US/10/580,660** TIME: 11:06:47

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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date